



A DOCPHOENIX

STIC INTERNAL

CRFD 9
Computer Readable Form Defective

TC INTERNAL

SEQREQ
Sequence Problem Att. From Exr.

CRFE
Computer Readable Form "ENTERED"

INTERNAL SEQUENCE DOCUMENT INDEX SHEET

STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING **ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: _____

Source: _____

Date Processed by STIC: _____

10/552,324
JFWO
12/20/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE **CHECKER VERSION 4.4.0 PROGRAM**, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

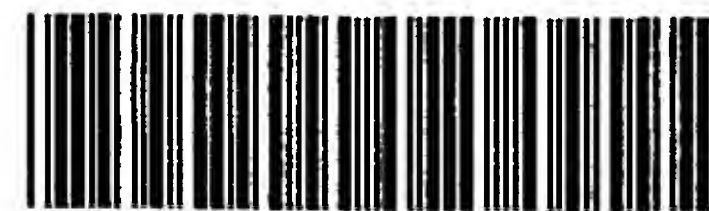
Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06



IFWO

RAW SEQUENCE LISTING

DATE: 12/20/2006

PATENT APPLICATION: US/10/552,324

TIME: 13:57:06

Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt

Output Set: N:\CRF4\12202006\J552324.raw

3 <110> APPLICANT: Igeneon Krebs-Immuntherapie Forschungs- & Entwickl
 5 <120> TITLE OF INVENTION: Immunogenic Recombinant Antibody
 7 <130> FILE REFERENCE: Immunogenic Recombinant AB
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/552,324
 C--> 10 <141> CURRENT FILING DATE: 2005-10-07
 12 <160> NUMBER OF SEQ ID NOS: 5
 14 <170> SOFTWARE: PatentIn Ver. 2.1
 16 <210> SEQ ID NO: 1
 17 <211> LENGTH: 3973
 18 <212> TYPE: DNA
 19 <213> ORGANISM: Artificial Sequence
 21 <220> FEATURE:
 22 <223> OTHER INFORMATION: Description of Artificial Sequence: mAB 17-1A
 24 <400> SEQUENCE: 1
 25 ataggctagc ctcgagccac caccatgcat cagaccagca tgggcatcaa gatggaatca 60
 26 cagactctgg tcttcatatc catactgctc tgggttatatg gagctgatgg gaacattgta 120
 27 atgacccaat ctcccaaate catgtccatg tcagtaggag agagggtcac cttgacctgc 180
 W--> 28 aaggccagtg agaattgtgt tacttatgtt tontgggtatc aacagaaacc agagcagtct 240
 29 cctaaactgc tgatatatgg ggcattccaa cggtagactg ggggtccnga tcgcttcaca 300
 30 ggcagtggat ctgcaacaga tttcactctg accatcagca gtgtgcaggc tgaagacctt 360
 31 gcagattatc actgtggaca gggttacagc tatccgtaca cgttcggagg ggggaccaag 420
 32 ctggaaataa aacgggctga tgctgcacca actgtatcca tcttcccacc atccagttag 480
 33 cagttaacat ctggaggtgc ctcatgctg tgcttcttga acaacttcta ccccaaagac 540
 34 atcaatgtca agtgaagat tgatggcagt gaacgacaaa atggcgtcct gaacagttgg 600
 35 actgatcagg acagcaaaga cagcacctac agcatgagca gcaccctcac gttgaccaag 660
 36 gacgagtatg aacgacataa cagctatacc tgtgaggcca ctcacaagac atcaacttca 720
 37 cccattgtca agagcttcaa caggaatgag tgtagacgc gtggatccgc ccctctccct 780
 38 ccccccccc taacgttact ggccgaagcc gcttggaata aggccggtgt gcgtttgtct 840
 39 atatgtgatt ttccaccata ttgccgtctt ttggcaatgt gagggcccgg aaacctggcc 900
 40 ctgtcttctt gacgagcatt cctaggggtc tttcccctct cgccaaagga atgcaaggct 960
 41 tgttgaatgt cgtgaaggaa gcagttcctc tggaagcttc ttgaagacaa acaactctct 1020
 42 tagcgaccct ttgcaggcag cggaaccccc cacctggcga caggtgcctc tgcggccaaa 1080
 43 agccacgtgt ataagataca cctgcaaagg cggcacaacc ccagtgccac gttgtgagtt 1140
 44 ggatagttgt ggaaagagtc aaatggctct cctcaagcgt attcaacaag gggctgaagg 1200
 45 atgcccagaa ggtaccccat tgtatgggat ctgatctggg gcctcgggtgc acatgcttta 1260
 46 catgtgttta gtcgagggtta aaaaaacgtc taggcccccc gaaccacggg gacgtggttt 1320
 47 tcctttgaaa aacacgatga taatatggcc accaccatgg aatggagcag agtctttatc 1380
 48 tttctcctat cagtaactgc aggtgttcac tcccaggctc agttgcagca gtctggagct 1440
 49 gagctggtta ggctgggac ttcagtgaag gtgtcctgca aggcttctgg atacgccttc 1500
 50 actaattact tgatagagtg ggtaaagcag aggcctggac agggccttga gtggattggg 1560
 51 gtgattaatc ctggaagtgg tggtaactaa tacaatgaga agttcaagg caaggcaaca 1620
 52 ctgactgcag acaaatcctc cagcactgcc tacatgcagc tcagcagcct gacatctgat 1680
 53 gactctgcgg tctatttctg tgcaagagat ggtccctggt ttgcttactg gggccaaggg 1740

Does Not Comply
 Corrected Diskette Needed
 CP8-1, 2, 4, 5
 Insufficient explanation
 Give the source
 of genetic
 material.

P/s Explain
 n' locations.
 See pg - 7
 for Error
 Explanation.

RAW SEQUENCE LISTING

DATE: 12/20/2006

PATENT APPLICATION: US/10/552,324

TIME: 13:57:06

Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt

Output Set: N:\CRF4\12202006\J552324.raw

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54 actctgggtca ctgtctctgc agccaaaaca acagcccccac cggtctatcc actggcccct 1800
55 gtgtgtggag atacaactgg ctccctcgggtg actctaggat gcctgggtcaa gggttatttc 1860
56 cctgagccag tgaccttgac ctggaactct ggatccctgt ccagtgggtgt gcacaccttc 1920
57 ccagctgtcc tgcagtctga cctctacacc ctcagcagct cagtgactgt aacctcgagc 1980
58 acctggccca gccagtccat cacctgcaat gtggcccacc cggcaagcag caccaagggtg 2040
59 gacaagaaaa ttgagcccag agggcccaca atcaagccct gtcctccatg caaatgccc 2100
60 gcacctaac tcttgggtgg accatccgtc ttcactcttc ctccaaagat caaggatgta 2160
61 ctcatgatct ccctgagccc catagtcaca tgtgtgggtg tggatgtgag cgaggatgac 2220
62 ccagatgtcc agatcagctg gtttgtgaac aacgtggaag tacacacagc tcagacacaa 2280
63 acccatagag aggattacaa cagtactctc cgggtgggtc gtgccctccc catccagcac 2340
64 caggactgga tgagtggcaa ggagttcaaa tgcaagggtc acaacaaaga cctcccagcg 2400
65 cccatcgaga gaacctctc aaaacccaaa gggtcagtaa gagctccaca ggtatatgtc 2460
66 ttgcctccac cagaagaaga gatgactaag aaacagggtc ctctgacctg catggtcaca 2520
67 gacttcatgc ctgaagacat ttacgtggag tggaccaaca acgggaaaac agagctaaac 2580
68 tacaagaaca ctgaaccagt cctggactct gatgggttctt acttcatgta cagcaagctg 2640
69 agagtggaaa agaagaactg ggtggaaaga aatagctact cctgttcagt ggtccacgag 2700
70 ggtctgcaca atcaccacac gactaagagc ttctcccgga ctccgggtaa atgagtcgac 2760
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74 agcattccta ggggtctttc ccctctcgcc aaaggaatgc aagggtctgtt gaatgtcgtg 3000
75 aaggaagcag ttctcttga agcttcttga agacaaacaa cgtctgtagc gacctttgc 3060
76 aggcagcgga accccccacc tggcgacagg tgcctctgcg gccaaaagcc acgtgtataa 3120
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83 cgctcaggaa cgagttcaag tacttccaaa gaatgaccac aacctcttca gtggaaggta 3540
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85 ctttaaagga cagaattaat atagttctca gtagagaact caaagaacca ccacgaggag 3660
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87 caagtaaagt agacatggtt tggatagtcg gaggcagttc tgtttaccag gaagccatga 3780
88 atcaaccagg ccacctcaga ctctttgtga caaggatcat gcaggaattt gaaagtgaca 3840
89 cgtttttccc agaaattgat ttggggaaat ataaacttct cccagaatac ccaggcgtcc 3900
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91 actaagcggc cgc 3973

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93 <210> SEQ ID NO: 2

94 <211> LENGTH: 465

95 <212> TYPE: PRT

96 <213> ORGANISM: Artificial Sequence

98 <220> FEATURE:

99 <223> OTHER INFORMATION: Description of Artificial Sequence: mAB 17-1A

101 <400> SEQUENCE: 2

102 Met Glu Trp Ser Arg Val Phe Ile Phe Leu Leu Ser Val Thr Ala Gly

103 1 5 10 15

105 Val His Ser Gln Val Gln Leu Gln Gln Ser Gly Ala Glu Leu Val Arg

106 20 25 30

Same Error

RAW SEQUENCE LISTING

DATE: 12/20/2006

PATENT APPLICATION: US/10/552,324

TIME: 13:57:06

Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt

Output Set: N:\CRF4\12202006\J552324.raw

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108 Pro Gly Thr Ser Val Lys Val Ser Cys Lys Ala Ser Gly Tyr Ala Phe
109          35          40          45
111 Thr Asn Tyr Leu Ile Glu Trp Val Lys Gln Arg Pro Gly Gln Gly Leu
112          50          55          60
114 Glu Trp Ile Gly Val Ile Asn Pro Gly Ser Gly Gly Thr Asn Tyr Asn
115 65          70          75          80
117 Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Lys Ser Ser Ser
118          85          90          95
120 Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Asp Asp Ser Ala Val
121          100          105          110
123 Tyr Phe Cys Ala Arg Asp Gly Pro Trp Phe Ala Tyr Trp Gly Gln Gly
124          115          120          125
126 Thr Leu Val Thr Val Ser Ala Ala Lys Thr Thr Ala Pro Ser Val Tyr
127          130          135          140
129 Pro Leu Ala Pro Val Cys Gly Asp Thr Thr Gly Ser Ser Val Thr Leu
130 145          150          155          160
132 Gly Cys Leu Val Lys Gly Tyr Phe Pro Glu Pro Val Thr Leu Thr Trp
133          165          170          175
135 Asn Ser Gly Ser Leu Ser Ser Gly Val His Thr Phe Pro Ala Val Leu
136          180          185          190
138 Gln Ser Asp Leu Tyr Thr Leu Ser Ser Ser Val Thr Val Thr Ser Ser
139          195          200          205
141 Thr Trp Pro Ser Gln Ser Ile Thr Cys Asn Val Ala His Pro Ala Ser
142          210          215          220
144 Ser Thr Lys Val Asp Lys Lys Ile Glu Pro Arg Gly Pro Thr Ile Lys
145 225          230          235          240
147 Pro Cys Pro Pro Cys Lys Cys Pro Ala Pro Asn Leu Leu Gly Gly Pro
148          245          250          255
150 Ser Val Phe Ile Phe Pro Pro Lys Ile Lys Asp Val Leu Met Ile Ser
151          260          265          270
153 Leu Ser Pro Ile Val Thr Cys Val Val Val Asp Val Ser Glu Asp Asp
154          275          280          285
156 Pro Asp Val Gln Ile Ser Trp Phe Val Asn Asn Val Glu Val His Thr
157          290          295          300
159 Ala Gln Thr Gln Thr His Arg Glu Asp Tyr Asn Ser Thr Leu Arg Val
160 305          310          315          320
162 Val Ser Ala Leu Pro Ile Gln His Gln Asp Trp Met Ser Gly Lys Glu
163          325          330          335
165 Phe Lys Cys Lys Val Asn Asn Lys Asp Leu Pro Ala Pro Ile Glu Arg
166          340          345          350
168 Thr Ile Ser Lys Pro Lys Gly Ser Val Arg Ala Pro Gln Val Tyr Val
169          355          360          365
171 Leu Pro Pro Pro Glu Glu Glu Met Thr Lys Lys Gln Val Thr Leu Thr
172          370          375          380
174 Cys Met Val Thr Asp Phe Met Pro Glu Asp Ile Tyr Val Glu Trp Thr
175 385          390          395          400
177 Asn Asn Gly Lys Thr Glu Leu Asn Tyr Lys Asn Thr Glu Pro Val Leu
178          405          410          415
180 Asp Ser Asp Gly Ser Tyr Phe Met Tyr Ser Lys Leu Arg Val Glu Lys

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RAW SEQUENCE LISTING

DATE: 12/20/2006

PATENT APPLICATION: US/10/552,324

TIME: 13:57:06

Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt

Output Set: N:\CRF4\12202006\J552324.raw

181 420 425 430
 183 Lys Asn Trp Val Glu Arg Asn Ser Tyr Ser Cys Ser Val Val His Glu
 184 435 440 445
 186 Gly Leu His Asn His His Thr Thr Lys Ser Phe Ser Arg Thr Pro Gly
 187 450 455 460

189 Lys

190 465

193 <210> SEQ ID NO: 3

194 <211> LENGTH: 243

195 <212> TYPE: PRT

196 <213> ORGANISM: Artificial Sequence

198 <220> FEATURE:

199 <223> OTHER INFORMATION: Description of Artificial Sequence (mAB 17-1A)

201 <400> SEQUENCE: 3

202 Met His Gln Thr Ser Met Gly Ile Lys Met Glu Ser Gln Thr Leu Val

203 1 5 10 15

205 Phe Ile Ser Ile Leu Leu Trp Leu Tyr Gly Ala Asp Gly Asn Ile Val

206 20 25 30

208 Met Thr Gln Ser Pro Lys Ser Met Ser Met Ser Val Gly Glu Arg Val

209 35 40 45

211 Thr Leu Thr Cys Lys Ala Ser Glu Asn Val Val Thr Tyr Val Ser Trp

212 50 55 60

214 Tyr Gln Gln Lys Pro Glu Gln Ser Pro Lys Leu Leu Ile Tyr Gly Ala

215 65 70 75 80

217 Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser

218 85 90 95

220 Ala Thr Asp Phe Thr Leu Thr Ile Ser Ser Val Gln Ala Glu Asp Leu

221 100 105 110

223 Ala Asp Tyr His Cys Gly Gln Gly Tyr Ser Tyr Pro Tyr Thr Phe Gly

224 115 120 125

226 Gly Gly Thr Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala Pro Thr Val

227 130 135 140

229 Ser Ile Phe Pro Pro Ser Ser Glu Gln Leu Thr Ser Gly Gly Ala Ser

230 145 150 155 160

232 Val Val Cys Phe Leu Asn Asn Phe Tyr Pro Lys Asp Ile Asn Val Lys

233 165 170 175

235 Trp Lys Ile Asp Gly Ser Glu Arg Gln Asn Gly Val Leu Asn Ser Trp

236 180 185 190

238 Thr Asp Gln Asp Ser Lys Asp Ser Thr Tyr Ser Met Ser Ser Thr Leu

239 195 200 205

241 Thr Leu Thr Lys Asp Glu Tyr Glu Arg His Asn Ser Tyr Thr Cys Glu

242 210 215 220

244 Ala Thr His Lys Thr Ser Thr Ser Pro Ile Val Lys Ser Phe Asn Arg

245 225 230 235 240

247 Asn Glu Cys

251 <210> SEQ ID NO: 4

252 <211> LENGTH: 243

253 <212> TYPE: PRT

254 <213> ORGANISM: Artificial Sequence

7 Same Error

RAW SEQUENCE LISTING

DATE: 12/20/2006

PATENT APPLICATION: US/10/552,324

TIME: 13:57:06

Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt

Output Set: N:\CRF4\12202006\J552324.raw

256 <220> FEATURE:

257 <223> OTHER INFORMATION: Description of Artificial Sequence:mAB 17-1A

259 <400> SEQUENCE: 4

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260 Met His Gln Thr Ser Met Gly Ile Lys Met Glu Ser Gln Thr Leu Val
261   1           5           10           15
263 Phe Ile Ser Ile Leu Leu Trp Leu Tyr Gly Ala Asp Gly Asn Ile Val
264           20           25           30
266 Met Thr Gln Ser Pro Lys Ser Met Ser Met Ser Val Gly Glu Arg Val
267           35           40           45
269 Thr Leu Thr Cys Lys Ala Ser Glu Asn Val Val Thr Tyr Val Ser Trp
270           50           55           60
272 Tyr Gln Gln Lys Pro Glu Gln Ser Pro Lys Leu Leu Ile Tyr Gly Ala
273   65           70           75           80
275 Ser Asn Arg Tyr Thr Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser
276           85           90           95
278 Ala Thr Asp Phe Thr Leu Thr Ile Ser Ser Val Gln Ala Glu Asp Leu
279           100          105          110
281 Ala Asp Tyr His Cys Gly Gln Gly Tyr Ser Tyr Pro Tyr Thr Phe Gly
282           115          120          125
284 Gly Gly Thr Lys Leu Glu Ile Arg Arg Ala Asp Ala Ala Pro Thr Val
285           130          135          140
287 Ser Ile Phe Pro Pro Ser Ser Glu Gln Leu Thr Ser Gly Gly Ala Ser
288   145          150          155          160
290 Val Val Cys Phe Leu Asn Asn Phe Tyr Pro Lys Asp Ile Asn Val Lys
291           165          170          175
293 Trp Lys Ile Asp Gly Ser Glu Arg Gln Asn Gly Val Leu Asn Ser Trp
294           180          185          190
296 Thr Asp Gln Asp Ser Lys Asp Ser Thr Tyr Ser Met Ser Ser Thr Leu
297           195          200          205
299 Thr Leu Thr Lys Asp Glu Tyr Glu Arg His Asn Ser Tyr Thr Cys Glu
300           210          215          220
302 Ala Thr His Lys Thr Ser Thr Ser Pro Ile Val Lys Ser Phe Asn Arg
303   225          230          235          240
305 Asn Glu Cys

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309 <210> SEQ ID NO: 5

310 <211> LENGTH: 243

311 <212> TYPE: PRT

312 <213> ORGANISM: Artificial Sequence

314 <220> FEATURE:

315 <223> OTHER INFORMATION: Description of Artificial Sequence:mAB 17-1A

317 <400> SEQUENCE: 5

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318 Met His Gln Thr Ser Met Gly Ile Arg Met Glu Ser Gln Thr Leu Val
319   1           5           10           15
321 Phe Ile Ser Ile Leu Leu Trp Leu Tyr Gly Ala Asp Gly Asn Ile Val
322           20           25           30
324 Met Thr Gln Ser Pro Arg Ser Met Ser Met Ser Val Gly Glu Arg Val
325           35           40           45
327 Thr Leu Thr Cys Arg Ala Ser Glu Asn Val Val Thr Tyr Val Ser Trp
328           50           55           60

```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/552,324

DATE: 12/20/2006
TIME: 13:57:07

Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt
Output Set: N:\CRF4\12202006\J552324_raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 213,288

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:5; Line(s) 368,369,370,371,372,373,374,375,376,377,378,379,380,381,382

VARIABLE LOCATION SUMMARY

DATE: 12/20/2006

PATENT APPLICATION: US/10/552,324

TIME: 13:57:07

Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt

Output Set: N:\CRF4\12202006\J552324.raw

Use of n's or Xaa's (NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.

Use of <220> to <223> is MANDATORY if n's or Xaa's are present.

in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Error Explanation

Seq#:1; N Pos. 213,288

VERIFICATION SUMMARY

DATE: 12/20/2006

PATENT APPLICATION: US/10/552,324

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Input Set : N:\efs\12_20_06\10552324_efs\10552324.txt

Output Set: N:\CRF4\12202006\J552324.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:28 M:258 W: Mandatory Feature missing, <221> Tag not found for SEQ ID#:1
L:28 M:258 W: Mandatory Feature missing, <222> Tag not found for SEQ ID#:1
L:28 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:180
M:341 Repeated in SeqNo=1